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OM nucleic - nucleic search, using sw mode

Run on: February 10, 2006, 05:26:42 ; Search time 4815 seconds

(without alignments)  
10270.779 Million cell updates/sec

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Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 5883141 seqs, 28421725653 residues

Total number of hits satisfying chosen parameters: 11766282

Minimum DB seq length: 0  
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : GenBank:  
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3: gb\_env::\*  
4: gb\_on::\*  
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9: gb\_ro::\*  
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13: gb\_vl::\*  
14: gb\_htg::\*  
15: gb\_pl::\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Result No.	Score	Query Match Length	DB ID	Description
1	870	100.0	AY293286	AY293286
2	857.2	98.5	CQ947366	AY293286 Homo sapiens B and T lymphocyte attenuator (BTLA) mRNA, complete cDNA sequence
3	857.2	98.5	CQ947428	AY293286.1 GI:31880026
4	854	98.2	CS10555	PR1 18-JUN-2002
5	845	97.2	AKL31204	mRNA, complete
6	764.2	87.8	CQ947365	linear mRNA, complete
7	704.2	80.9	CQ947364	PR1 18-JUN-2002
8	527.6	60.6	CQ947360	mRNA, complete
9	520.2	59.8	BDD00983	PR1 18-JUN-2002
10	466.2	53.6	CQ947358	mRNA, complete
11	437.6	50.3	CQ947359	PR1 18-JUN-2002
12	425	48.9	CS03517	mRNA, complete
13	425	48.9	CS04469	mRNA, complete
14	402.8	46.3	BD13124	mRNA, complete
15	402.8	46.3	CS13203	mRNA, complete
16	402.8	46.3	AR33978	mRNA, complete
17	389	44.6	CQ947363	mRNA, complete
18	378	43.4	BD131144	mRNA, complete

SUMMARIES

REFERENCE	AUTHORS	JOURNAL	TITLE	DESCRIPTION
1	(bases 1 to 870)	Nat. Immunol.	In press	Mammalia; Ruthenica; Euarchontoglires; Primates; Catarrhini; Homidae; Homo.
2	(bases 1 to 870)	Watanaabe,N., Gavrieli,M., Sedy,J.R., Yang,J., Fallarino,F., Loftin,S.K., Hurchla,M.A., Zimmerman,N., Sim,J., Zang,X., Murphy,T.L., Russell,J.H., Allison,J.P. and Murphy,K.M.	Submitted (08-MAY-2003) Pathology, Washington University, 660 S. Euclid, St. Louis, MO 63110, USA	BTLA is a lymphocyte inhibitory receptor with similarities to CTLA-4 and PD-1
3	Murphy,K.M., Watanaabe,N., Yang,J. and Murphy,T.L.	Nat. Immunol.	In press	BTLA is a lymphocyte inhibitory receptor with similarities to CTLA-4 and PD-1
4	Watanaabe,N., Gavrieli,M., Sedy,J.R., Yang,J., Fallarino,F., Loftin,S.K., Hurchla,M.A., Zimmerman,N., Sim,J., Zang,X., Murphy,T.L., Russell,J.H., Allison,J.P. and Murphy,K.M.	Submitted (08-MAY-2003) Pathology, Washington University, 660 S. Euclid, St. Louis, MO 63110, USA	BTLA is a lymphocyte inhibitory receptor with similarities to CTLA-4 and PD-1	
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ALIGNMENTS

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## ORIGIN

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 Matches 870; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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## RESULT 2

CQ947366 CQ947366 870 bp DNA linear PAT 01-DBCC-2004

LOCUS CQ947366 Sequence 42 from Patent WO2004096976.

DEFINITION CQ947366 Sequence 42 from Patent WO2004096976.

ACCESSION CQ947366.1 GI:56295403 Version 1

SOURCE Homo sapiens (human)

ORGANISM Homo sapiens (human)

Hominidae; Homo.

1 KAYE, J. and WILKINSON, B.

REFERENCE Spex compositions and methods of use

AUTHORS Spex compositions and methods of use

JOURNAL Patent: WO 2004096976-A 42 11-NOV-2004; Novartis AG (CH); Novartis Pharma GmbH (AT); The Scripps Research

PRATURES Institute (US)

Source Location/Qualifiers

1. .870

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/mol\_type="unassigned DNA"

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## ORIGIN

Query Match 98.5%; Score 857.2; DB 6; Length 870;  
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 Matches 862; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

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61 CCATATCTGGACATCTGGAAATCTGATGAGGAAAGATCTGATGAGCTTATA 120

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121 AAGAGAACATCTGAAACACTCTACATCTGAGGAGATCTGATGAGCTTATA 180

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241 GAAAGACCTCAAGGAGGAAGTGGAGAGCAAGACGCCCTGGCTCTGTTAGTTACTCT 300

301 CATTGAAACCATGCTCTCTATGACATGGCTACCCCTGTCATTCATTTCAG 360

301 CATTGAAACCATGCTCTCTATGACATGGCTACCCCTGTCATTCATTTCAG 360

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 Qy 301 CATTGAGACCATGCTCTTAAATGACATGGATCTGGTCAATTTCAG 360  
 Db 397 CATTGAGACCATGCTCTTAAATGACATGGATCTGGTCAATTTCAG 456  
 Qy 361 TCTAATCTCATGAAAGCCACTCACACTTTAATGACGTGATAAGGTCTTA 420  
 Db 457 TCTAATCTCATGAAAGCCACTCACACTTTAATGACGTGATAAGGTCTTA 516  
 Qy 421 GAAAGACCTCCAGGAGCAGACTGGCGAGAGACCCCTGGCTCTGATAGTTACTCT 480  
 Db 517 GAAAGACCTCCAGGAGCAGACTGGCGAGAGACCCCTGGCTCTGATAGTTACTCT 576  
 Qy 481 TGGGGGATGGCTCTACTCATCACTACCTGGTCTGCTCTGCTGCTGATAGTTACTCT 540  
 Db 577 TGGGGGATGGCTCTACTCATCACTACCTGGTCTGCTCTGCTGCTGATAGTTACTCT 636  
 Qy 541 CACTAGGAAAGCAAAATGACTCTAACAGCAGGAGGAATATCTGGTAT 600  
 Db 637 CACTAGGAAAGCAAAATGACTCTAACAGCAGGAGGAATATCTGGTAT 696  
 Qy 601 ACTCACTTAAGCAGGAGCAACAGAGACCAACCCAGGCAAATTCCAGTACTCTA 660  
 Db 697 OCTCACTTAAGCAGGAGCAACAGAGACCAACCCAGGCAAATTCCAGTACTCTA 756  
 Qy 661 TCAAACTGGATTATGATAATGACCTGACCTTGTGGAGGTCTGAGAAGGTCT 720  
 Db 757 TCAAACTGGATTATGATAATGACCTGACCTTGTGGAGGTCTGAGAAGGTCT 816  
 Qy 721 GAGTTGTTCTAATCCTGAAAGAACACAGGATTTATCTCCCTG 780  
 Db 817 GAGTTGTTCTAATCCTGAAAGAACACAGGATTTATCTCCCTG 876  
 Qy 781 AACATTCTGGTCTGGACTGAACTGAGCTGGAGAAGATGAAAGACACACA 840  
 Db 877 AACATTCTGGTCTGGACTGAACTGAGCTGGAGAAGATGAAAGACACACA 936  
 Qy 841 GAATATGATCCATGTTGGAGGTTAA 870  
 Db 937 GATATGATCCATGTTGGAGGTTAA 966

RESULT 7

US-10-371-341-1

/ Sequence 1, Application US/10371341  
 / Publication No. US20040091884A1  
 / GENERAL INFORMATION:  
 / APPLICANT: HILARY CLARK  
 / APPLICANT: DANIEL L. BATON  
 / APPLICANT: AUSTIN L. GURNEY  
 / APPLICANT: BERND WANK  
 / TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR THE TREATMENT OF  
 / TITLE OF INVENTION: IMMUNE RELATED DISORDERS  
 / FILE REFERENCE: P19561-US  
 / CURRENT APPLICATION NUMBER: US/10/371,341  
 / PRIOR FILING DATE: 2003-02-19  
 / PRIOR APPLICATION NUMBER: US 60/421,236  
 / PRIOR FILING DATE: 2002-10-25  
 / NUMBER OF SEQ ID NOS: 2  
 / SEQ ID NO: 1  
 / LENGTH: 1066  
 / TYPE: DNA  
 / ORGANISM: Homo sapien

Db 24 ATGAGACATGCCCTCCATCTCTGGACTGGAAATATTGGGCTCTTCATAC 83  
 Qy 61 CCATATCTGGACATCTGGACATCTGGAAAGATCATGATGAGTACAGTTATA 120  
 Db 84 CCATATCTGGACATCTGGACATCTGGAAAGATCATGATGAGTACAGTTATA 143  
 Qy 121 AAGAGCAAACTGACACTCATTTAGCAGGAGTCCTTGACTAGATGCCTG 180  
 Db 144 AAGAGCAAACTGACACTCATTTAGCAGGAGTCCTTGACTAGATGCCTG 203  
 Qy 181 AAATCTGTGTAAGGCTCATCTGACTGCTGCTGATAGTTACTCTA 240  
 Db 204 AAATCTGTGTAAGGCTCATCTGACTGCTGCTGATAGTTACTCTA 263  
 Qy 241 AACTGAGATGACAAACAGTGGCTGACTGGTGGCTGAGCTGATAGTTACTCT 300  
 Db 264 AACTGAGATGACAAACAGTGGCTGACTGGTGGCTGAGCTGATAGTTACTCT 323  
 Qy 301 CATTGAGACCATGCTCTAACAGCAGGAGGAATATCTGGTAT 360  
 Db 324 CATTGAGACCATGCTCTAACAGCAGGAGGAATATCTGGTAT 383  
 Qy 361 TCTAATCTCATGAAAGCAAAATGACTCTAACAGCAGGAGGAATATCTGGTAT 420  
 Db 384 TCTAATCTCATGAAAGCAAAATGACTCTAACAGCAGGAGGAATATCTGGTAT 443  
 Qy 421 GAAAGACCTCCAGGAGCAACAGGAGCAACCCAGGCAAATTCCAGTACTCTA 480  
 Db 444 GAAAGACCTCCAGGAGCAACAGGAGCAACCCAGGCAAATTCCAGTACTCTA 503  
 Qy 481 TGGGGGATGGCTCTACTCATCTACTGGTCTGCTGCTGCTGCTGCTGAGG 540  
 Db 504 TGGGGGATGGCTCTACTCATCTACTGGTCTGCTGCTGCTGCTGAGG 563  
 Qy 541 CACCAAGAAAGCAAAATGACTCTGACAGCAGGAGCAACCCAGGCAAATTCCAGTACTCTA 600  
 Db 564 CACCAAGAAAGCAAAATGACTCTGACAGCAGGAGCAACCCAGGCAAATTCCAGTACTCTA 623  
 Qy 601 GCTCACTTAAGGAGGAGGAGCAACAGGAGCAACCCAGGCAAATTCCAGTACTCTA 660  
 Db 624 GCTCACTTAAGGAGGAGGAGCAACAGGAGCAACCCAGGCAAATTCCAGTACTCTA 683  
 Qy 661 TCAGAGCTGGATTATGATAATCCTGACTTGTGGAGGAGGTCT 720  
 Db 684 TCAGAGCTGGATTATGATAATCCTGACTTGTGGAGGAGGTCT 743  
 Qy 721 GAGTTGTTCTAATCCTGAAAGAACACAGGATTTATCTCCCTG 780  
 Db 744 GAGTTGTTCTAATCCTGAAAGAACACAGGATTTATCTCCCTG 803  
 Qy 781 AACATTCTGGTCTGGACTGAACTGAGCTGGAGAAGATGAAAGACACACA 840  
 Db 804 AACATTCTGGTCTGGACTGAACTGAGCTGGAGAAGATGAAAGACACACA 863  
 Qy 841 GAATATGATCCATGTTGGAGGTTAA 870  
 Db 864 GAATATGATCCATGTTGGAGGTTAA 893

RESULT 8

US-10-989-926-27

/ Sequence 27, Application US/10989826  
 / Publication No. US20050238650A1  
 / GENERAL INFORMATION:  
 / APPLICANT: Generitech, Inc.  
 / APPLICANT: Crowley, Craig  
 / APPLICANT: De Savage, Frederic J.  
 / APPLICANT: Baton, Daniel L.  
 / APPLICANT: Ebens, Allen  
 / APPLICANT: Polson, Andrew  
 / APPLICANT: Smith, Victoria  
 / TITLE OF INVENTION: Compositions and Methods for the Treatment of Tumor of  
 / TITLE OF INVENTION: Hematopoietic Origin

Query Match 98.2%, Score 854, DB 7, length 1066;  
 Best Local Similarity 98.9%, Pred. No. 2, Bp-246;  
 Matches 860, Conservative 0, Mismatches 10; Indels 0, Gaps 0;

1 ATGAGACATGCCCTCCATCTCTGGACTGGAAATATTGGGCTCTTCATAC 60

Qy US-10-371-341-1



5

## RESULT 2

US-10-371-341-2

Sequence 2, Application US/10371341

## GENERAL INFORMATION:

APPLICANT: HILARY CLARK

APPLICANT: DANIEL L. BATON

APPLICANT: AUSTIN L. GRUNBY

APPLICANT: BERND WRAK

## TITLE OR INVENTION: NOVEL COMPOSITIONS AND METHODS FOR THE TREATMENT OF

FILE REFERENCE: P1956R1-US

CURRENT APPLICATION NUMBER: US/10/371,341

CURRENT FILING DATE: 2003-02-19

PRIORITY NUMBER: US 60/421,236

PRIOR FILING DATE: 2002-10-25

SEQ ID NO 2

LENGTH: 289

TYPE: PRT

ORGANISM: Homo sapien

US-10-371-341-2

Query Match 97.4%; Score 1516; DB 4; Length 289;  
 Best Local Similarity 97.9%; Pred. No. 1.5e-145; Indels 0; Gaps 0;  
 Matches 283; Conservative 2; Mismatches 4; Indels 0; Gaps 0;

Qy 1 MKTLPAMLGTKLFWVFLIPYLIPYDINWINGKESCDVOLYKROSEHSLAGDPPELCRV 60

Db 1 MKTLPAMLGTKLFWVFLIPYLIPYDINWINGKESCDVOLYKROSEHSLAGDPPELCRV 60

Qy 61 KYCANPRPHTWCKLNGTCVKLDRQTSWERKNSCSPTRIHPMLPNDNGSYRCANFO 120

Db 61 KYCANPRPHTWCKLNGTCVKLDRQTSWERKNSCSPTRIHPMLPNDNGSYRCANFO 120

Qy 121 SNLJSHSTLYTDVGASERPSKDEVASREWLLSPLPGGLPLITTCUCCAR 180

Db 121 SNLJSHSTLYTDVGASERPSKDEVASREWLLSPLPGGLPLITTCUCCAR 180

Qy 181 HQGKNEELDTAGREINLDAHLKSBOEASTRONSQVLSEAGYDNDPDCPRMORG 240

Db 181 HQGKNEELDTAGREINLDAHLKSBOEASTRONSQVLSEAGYDNDPDCPRMORG 240

Qy 181 HQGKNEELDTAGREINLDAHLKSBOEASTRONSQVLSEAGYDNDPDCPRMORG 240

Db 181 HQGKNEELDTAGREINLDAHLKSBOEASTRONSQVLSEAGYDNDPDCPRMORG 240

Qy 241 EVCSNPCLRENPGIVVASYLNSRARNVKAPEVASICVRS 289

Db 241 EVCSNPCLRENPGIVVASYLNSRARNVKAPEVASICVRS 289

Qy 241 EVCSNPCLRENPGIVVASYLNSRARNVKAPEVASICVRS 289

Db 241 EVCSNPCLRENPGIVVASYLNSRARNVKAPEVASICVRS 289

RESULT 4

US-10-331-622-21

Sequence 21, Application US/10831622

Publication No. US20040248257A1

## GENERAL INFORMATION:

APPLICANT: KAYE, Jonathan

APPLICANT: Wilkinson, Beverley

## TITLE OR INVENTION: SPEX COMPOSITIONS AND METHODS OF USE

FILE REFERENCE: T101 8.0.1

CURRENT APPLICATION NUMBER: US/10/831,622

CURRENT FILING DATE: 2004-04-23

PRIORITY NUMBER: US 60/467,206

PRIOR FILING DATE: 2003-04-30

NUMBER OF SEQ ID NOS: 113

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 21

LENGTH: 289

TYPE: PRT

ORGANISM: Homo sapiens

US-10-600-997-6

Sequence 6, Application US/10600997

Publication No. US20040175380A1

## GENERAL INFORMATION:

APPLICANT: Allison, James

APPLICANT: Murphy, Kenneth

APPLICANT: Watanabe, Norihiro

APPLICANT: Murphy, Theresa

APPLICANT: Yang, Jianfei

APPLICANT: Zang, Jingxing

## TITLE OR INVENTION: Compositions and Methods for Modulating Lymphocyte Activity

FILE REFERENCE: A-71608/TM/DDR

CURRENT APPLICATION NUMBER: US/10/600,997

CURRENT FILING DATE: 2003-06-20

PRIOR APPLICATION NUMBER: US 60/390,653

PRIOR FILING DATE: 2002-06-20

NUMBER OF SEQ ID NOS: 56

SOFTWARE: PatentIn version 3.2

SEQ ID NO 6

LENGTH: 289

Query Match 97.4%; Score 1516; DB 5; Length 289;

Best Local Similarity 97.9%; Pred. No. 1.5e-145; Indels 0; Gaps 0;

Matches 283; Conservative 2; Mismatches 4; Indels 0; Gaps 0;

Qy 1 MKTLPAMLGTKLFWVFLIPYLIPYDINWINGKESCDVOLYKROSEHSLAGDPPELCRV 60

Db 1 MKTLPAMLGTKLFWVFLIPYLIPYDINWINGKESCDVOLYKROSEHSLAGDPPELCRV 60

Qy 61 KYCANPRPHTWCKLNGTCVKLDRQTSWERKNSCSPTRIHPMLPNDNGSYRCANFO 120

Db 61 KYCANPRPHTWCKLNGTCVKLDRQTSWERKNSCSPTRIHPMLPNDNGSYRCANFO 120

Qy 121 SNLJSHSTLYTDVGASERPSKDEVASREWLLSPLPGGLPLITTCUCCAR 180

Db 121 SNLJSHSTLYTDVGASERPSKDEVASREWLLSPLPGGLPLITTCUCCAR 180

Qy 181 HQGKNEELDTAGREINLDAHLKSBOEASTRONSQVLSEAGYDNDPDCPRMORG 240

Db 181 HQGKNEELDTAGREINLDAHLKSBOEASTRONSQVLSEAGYDNDPDCPRMORG 240

Qy 241 EVCSNPCLRENPGIVVASYLNSRARNVKAPEVASICVRS 289

Db 241 EVCSNPCLRENPGIVVASYLNSRARNVKAPEVASICVRS 289